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MPsrch_pp protein - protein database search, using Smith-Waterman algorithm

Run on: Sat May 13 08:40:18 2000; MasPar time 4.47 Seconds 318.123 Million cell updates/sec

Tabular output not generated.

Title:

Description:
Perfect Score:
Sequence:

>US-09-331-631-21 (32-91) from US09331631.pep 453 1 TENPCAQRCIQSCQQEPDDL......DTGATNQRHPPGERTRGRQP 60

Scoring table: PAM 150 Gap 11

Searched: 188963 seqs, 23686106 residues

Post-processing: Minimum Match 0% Listing first 45 summaries

Database: a-geneseq35 1:geneseqp

Statistics: Mean 24.616; Variance 91.839; scale 0.268

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

18 70 15.5 19 70 15.5 20 70 15.5 21 70 15.5	70 15 70 15 70 15	70 15 70 15 70 15	70 15 70 15	70 15		72 15	73 16.		75 16	75	77 17			81 17	17	83 18	20	5 93 20.5	26		<u>سر</u>	1 453 100.0	Score Match	Result Ouerv
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T VIIIOCVIE MEMORANE	D Timoboonto monbusto	Soluble human insulin	Rat thrombomodulin.	Partial rat thrombomod	Partial rat thrombomod	H. pylori membrane pro	Zea mays antimicrobial	Novel human protein, d	Protein cognate of pro	Human insulin receptor	Macadamia integrifolia		Gossypium hirsutum ant	Sequence encoded by 67		Macadamia integrifolia	Sequence encoded by a	Human KAL protein.	Glycine max antimicrob	Peanut allergen Ara hI	Peanut allergen Ara hI	Arachis hypogaea antim	Description	
/. Z¤e+UL	7	7.286+01	7.28e+01	7.28e+01	7.28e+01	4.94e+01	4.06e+0i	3.34e+01	2.74e+01	2.74e+01	1.84e+01	1.84e+01	1.84e+01	8.24e+00	8.24e+00		6.89e-01	6.89e-01	2.44e-03	1.02e-32	4.98e-39	4.98e-39	Pred. No.	

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A desaturase enzyme en	Human desaturase enzym	Human cyclin D1/cyclin	Human cyclin D1-human	Amino acid sequence of	Human desaturase enzym	A desaturase enzyme en	Amino acid sequence of	H4-1BB receptor protei	Human 4-1BB receptor.	Maleate isomerase sequ	Human h4-1BBSV recepto	Human h4-1BBSV recepto	Human receptor induced	HSV-2 strain SB5 Conti	Poxvirus ORF-2 (VEGF r	Poxvirus ORF-2 protein	Poxvirus ORF-2 truncat	Poxvirus ORF-2 truncat	Poxvirus ORF-2 truncat	Poxvirus ORF-2 truncat	Wild type human insuli
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ALIGNMENTS

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<u> </u>	/label= Mat_protein	4		Peptide 1.22	cnrs nypogaea	monoclonal antibody; ELISA; analysis; Ara hI.	anaphylactic shock; immunotherapy; thera	; seed storage	gen Ara	29-DEC-1997 (first entry)		22110 5+5545+4.	32 TENPCAQRCLQSCQQEPDDLKQKACESRCTKLEYDPRCVYDTGATNQRHPPGERTRGRQP 91	32 TENPCAQRCLQSCQQEPDDLKQKACESRCTKLEYDPRCVYDTGATNQRHPPGERTRGRQP 91	Query Match 100.0%; Score 453; DB 1; Length 614; Best Local Similarity 100.0%; Pred. No. 4.98e-39; Matches 60; Conservative 0; Mismatches 0; Indels 0; Gaps	Sequence 614 AA;	imals.	used to control microbial infestations in plants and	nce is that of an antimicrobial	1: Page 55-57: 96pp: English.	l for controlling microbial infest	l anti-microbial protein from e o Macadamia	BOWER NI, GOUITER KC, Green JL, Manners JM, Marcus JP; WDT: 98-377279/32	OOP RES CENT TROPICAL PLANT PATHOLOGY.	20-DEC-1996; AU-004275.	22-DEC-1997; AU0874.	02-JUL-1998.	W09827805-A1.	(4717)	icrobial prote	1998 (first ent		US 1 W62834 standard: Dentide: 614 AA

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RESULT

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C This polypeptide comprises major peanut allergen Ara HI (W22149).

C Its sequence was deduced from cDNA clone P17 (T76612), isolated

C from peanut seed cDNA using a primer (see T76616) based on an

C isolated Ara HI peptide (see W24206). The sequence shows

c significant homology with the vicilin family of seed storage

groteins of other legumes. The allergen is recognised by serum

C proteins of other legumes. The allergen is recognised by serum

C IgE from a large proportion of individuals with peanut

C hypersensitivity. Ara HI and Ara HII (see W24164) can be used to

C raise monoclonal antibodies which are used in a specific two-site

C MAD ELISA for the detection of Ara HI or Ara HII (claimed). Ige-

C binding Ara HI antigen epitopes (see W24165-87) may be used in

C vaccines to protect against allergic reactions to peanut allergens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local Similarity 100.0%;
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10-JUL-1997.
23-SEP-1996; U15222.
04-MAR-1996; US-610424.
29-DEC-1995; US-009455.
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10-JUL-1997.
23-SEP-1996; U15222.
04-MAR-1996; US-610424.
29-DEC-1995; US-009455.
(UYAR-) UNIV ARKANSAS.
(UYAR-) UNIV ARKANSAS.
This polypeptide comprises major peanut allergen Ara hI (W22149). Its sequence was deduced from cDNA clone P41b (T76613), isolated from peanut seed cDNA using a primer (see T76616) based on an isolated Ara hI peptide (see W24206). The sequence shows significant homology with the vicilin family of seed storage proteins of other legumes. The allergen is recognised by serum IgE from a large proportion of individuals with peanut hypersensitivity. Ara hI and Ara hII (see W24164) can be used to raise monoclonal antibodies which are used in a specific two-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peanut; seed storage protein; allergen; allergy; hypersensitivity; vaccine; anaphylactic shock; immunotherapy; therapy; monocional antibody; ELISA; analysis; Ara hI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Arachis hypogaea strain Florunner.
Key Location/Qualifiers
                                                                                                                                                                                                                                                                        two-site monoclonal antibody based Claim 31; Page 172; 354pp; English.
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Bannon GA, Burks AW,
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Pred. No. 4.98e-39;
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W09824898-A2.
11-JUN-1998.
05-DEC-1997;
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22-DEC-1997;
20-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful for controlling microbial infestations of plants Claim 1; Page 63-65; 96pp; English.
The sequence is that of an antimicrobial protein which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY. Bower NI, Goulter KC, Green JL, Manners JM, Mar WPI; 98-377279/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         W62838 standard; Protein; W62838; (first entry) Glycine max antimicrobial
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                                                                                                                                                                                    Domain '
                                                                                                                                                                                                                                                               Peptide
                                                                                                                                                                                                                                                                                                                                                                                                     Human KAL protein: retinal
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                                                                                                                                                                                                                                                                                                                                                                               injury;
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Similarity 32.2%;
19; Conservative
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  È06806
                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                  fibronectin
                                                                                                                                             403.
                                                                                                                                                                                                                                                                                                           Location/Qualifiers 181. .285
                                                                                                                                                                                         286. .402
                                                                      /note=
                                                                                                                    /note=
                                                                                                                                                                   /note=
                                                                                                                                                                                                                                  /note= "32R1
                                                                                                                                                                                                                                                               254. .285
                                                                                                                                                                                                                                                                                     /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein;
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90.6%;
                                                                                               . 661
                                                                                                                                             . 540
                                                                                                                    "Fibronectin
                                                                                                                                                                "Fibronectin
                                                                                                                                                                                                                                                                                   "Fibronectin type III repeat"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     infestation;
                                                                      "Fibronectin
                                                                                                                                                                                                               32R1 peptide; this inventors under cla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein
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                                                                                                                                                                                                                                                                                                                                                                             injury; renal injury; neuronal
type III repeat; 32R1 peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 119; DB 1; Le Pred. No. 2.44e-03; 11; Mismatches 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score
Pred.
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pial infestations of
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No. 1.02e-32;
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claim 2 in the
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                                                                      III repeat"
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Matches 1
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Best Local Similarity
Matches 13; Conser
                                                                                                                                                                                                                                                 N-PSDB; Q40341.

Nucleic acid sequence of gene with X-linked Kallmann syndrome useful for diagnosing Kallmann syndrome by amplification to detect genetic anomalies

Disclosure; Fig 6; 60pp; French.

The nucleic acid sequence is derived from the ADMLX gene associated with KS (or Hypogonadotrophic hypogonadism and anosmia).

Oligonucleotide pairs which act as primers for specific amplification of the gene are used in amplification methods to detect genetic anomalies which cause KS. The primer pairs corresp. to the coding and non-coding regions of exon 1 of the ADMLX gene and one pair each for the other 13 exons.

Sequence 680 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LT 6
R34445 standard; Protein;
R34445;
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The present sequence represents the human KAL protein. The KAL gene is involved in morphogenetic events in neurons and kidney cells. The invention claims for a therapeutic composition comprising the KAL protein or one of its biologically active derivatives, e.g. the fibronectin type III repeats and the 32R1 peptide. This therapeutic composition is claimed to be useful for the treatment of retinal, remembers and neural injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-OCT-1992; F00956.
09-OCT-1991; FR-012451.
09-OCT-1991; FR-012451.
(INSP) INST PASTEUR.
(USSH) US DEPT HEALTH & HUMAN SERVICE.
Claverie J, Hardelin J, Legouis R, L
Lutfalla G, Petit C;
WPI: 93-134456/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-AUG-1993 (first entry)
09-AUG-1993 (first entry)
Sequence encoded by a cDNA clone contg. the ADMLX gene associated with x-linked Kallmann Syndrome (KS).
x-linked Kallmann syndrome; ADMLX gene; diagnosis; PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ardouin O, Hardelin J, Legouis R, Mazie J, Petit C, Rougon G, Sarailh C, Soussi-Yanicostas N;
WPI; 98-33318/29
Use of KAL protein - for the treatment of traumatic, infectious, metabolic or inherited nerve, spinal, retinal or renal injury or
                   W62830;
27-OCT-1998
Macadamia integrifolia antimicrobial protein
                                                  W62830 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9307267-A.
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(CNRS ) CNRS CENT NAT RECH
(INSP ) INST PASTEUR.
Ardouin O, Hardelin J, Lego
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Local Similarity 33.3%;
nes 13; Conservative
                                                                                                                  34 NPCAQRCLQSCQQEPDDLKQKACESRCTKLEYDPRCVYD
                                                                                                                                     81 KQCS-KCLEPCK-ESGDLRKHQCQSFCEPL-F-PKKSYE 115 : |: :||: |: |: |: |: |:
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ilarity 33.3%;
Conservative
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Pred.
11; M
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Pred. No. 6.89e-01
11; Mismatches 1
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1. No. 6.89e-01;
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Best Local S
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Matches 10; Conse
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22-DEC-1997; AU0874.

20-DEC-1996; AU-004275.

(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
Bower NI, Goulter KC, Green JL, Manners JM, Marc
WPI; 98-377279/32.

Novel anti-microbial protein from e.g. Macadamia
useful for controlling microbial infestations of
claim 1; Page 47-49; 96pp; English.
The sequence is that of an antimicrobial protein
be used to control microbial infestations in plan
R20181; standard; Protein; 1
R20181;
16-APR-1992 (first entry);
Sequence encoded by 67 kD:
Cocoa; flavour; vicilin; so
Theobroma cacao.
W09119801-A.
26-DEC-1991.
07-JUN-1991; G00914.
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(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
Bower NI, Goulter KC, Green JL, Manners JM, Marcus
WPI; 98-377279/32.

N-PSDB; V42316.
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W62831;
27-OCT-1998 (fin
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The sequence is that of an antimicrobial protein which can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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QRCLQSCQQEP-DDLKQKACESRCTKLEY
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Similarity 37.5%;
12; Conservative
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larity 34.5%;
Conservative
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67 kD T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "mature protein"
                                                                                                                                      seed storage protein.
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Pred.
8; M
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Pred. No. 5.48e
6; Mismatches
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5.48e+00;
5.2 12;
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stations of
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(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.

Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;

MPI; 98-377279/32.
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Claim 4; Fig 2; 59pp; English.

The inventors claim a 67 kD and 31 kD T. cacao protein, and fragments, and encoding DNAs. The 47 kD and 31 kD proteins are derived from the 67 kD precursor. T. cacao protein CDNA was detected in a cDNA library prepared from immature cocoa beans RNA using a probe based on the AA sequence of a CNBr peptide common to the 47 kD and 31 kD polypeptides. Homology searches revealed close homologies between the 67 kD polypeptide and the vicilins, which are
   WO9827805-A1.
02-JUL-1998.
                                                                                                            Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel anti-microbial protein from e.g. Macadamia integrifolia useful for controlling microbial infestations of plants or man Claim 1; Page 49-51; 96pp; English.

The sequence is that of an antimicrobial protein which can
                                                                                                                                                                   Peptide
                                                                                                                                                                                                                  Key
                                                                                                                                                                                                                                                      Macadamia integrifolia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 be used to control microbial infestations animals.
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22-DEC-1997; AU0874.
20-DEC-1996; AU-0042
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W62832 standard;
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(MRSC ) MARS UK LTD.
Spencer ME, Hodge R, Deakin
WPI; 92-024418/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Recombinant cocoa proteins -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         90
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Local Similarity 34.5%;
les 10; Conservation
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ence 566 AA;
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29. .666
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39.1%;
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yeast and bacterial
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25-NOV-1987.
19-MAY-1987; 200940.
20-MAY-1986; NL-001271.
(UYKA-) Katholieke Univ.
Van de Ven WJM,
                                                                                   Furin; insulin receptor; tumor
                                                                                                               T7 13

P70057 standard; protein; 149

P70057;

27-JAN-1991 (first entry)
                                                                                                                                                                                                                                                                                                be used to control microbial infestations animals.
                                                                Homo sapiens.
EP-246709-A.
                                                                                                    Human insulin receptor
                                                                                                                                                                                                                                                                                                                     useful for controlling microbial infestations of plants Claim 1; Page 39-41; 96pp; English.
The sequence is that of an antimicrobial protein which
                                                                                                                                                                                                                                                                                                                                                                             02-JUL-1998.
22-DEC-1997; AU0874.
22-DEC-1996; AU-004275.
20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
Bower NI, Goulter KC, Green JL, Manners JM, Mar
MPI; 98-377279/32.
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Macadamia integrifolia antimicrobial protein.
antimicrobial protein; infestation; control.
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W62829;
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The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and management.
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20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
Bower NI, Goulter KC, Green JL, Manners JM, Mar
WPI; 98-377279/32.
                                                                                                                                                                                                                                                                                     Sequence
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11; Conser
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Pred. No. 1.
6; Mismatc
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plants or mamma
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SCC CCC CCC PTT PTI
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                                                                                                This sequence is a protein cognate of protein kinase C-theta (PKC-C) theta) that binds to theta VI but not to delta VI of Iamin. The amino acid sequence was deduced from cDNA clone #10 (760588).

C amino acid sequence was deduced from cDNA clone #10 (760588).

C isolated from a human CD4+ T-cell DNA library using a yeast two-cylorid system. A claimed method for prepn. of the cognate involves culturing recombinant host cells under conditions where the cognate is produced. The cognate is used in a claimed method for identifying modulators of intracellular signal transduction. This involves cassessing the ability of candidate modulators to affect the interaction between a catalytically active signal generating protein (e.g. PKC-theta) and a cognate binding protein involved in modulating the signal transduction function. Substances identified using this method are used to modulate the immune system in a subject (claimed). They act to reduce T-cell activity, reduce the rate of graft rejection, reduce the severity of an autoimmune disorder, ameliorate an allergic and/or asthmatic response, or diminish cytokine response
 Query Match
Best Local Similarity
Matches 10; Conser
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kauvar LM, Mochly-Rosen D, Vasquez NJ, Voronova A; WPI; 97-236030/21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Protein cognate of protein kinase C-theta (clone #10 product). Signal transduction; cell signalling; modulator; immunomodulator; human; protein kinase C-theta; PKC-theta; cognate; graft rejection; autoimmune disease; allergy; asthma; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Fig 10; 24pp; English.

The sequence of the human insulin receptor is provided for comparison with the cysteine-rich region of furin. Furin is strongly expressed in specific types of tumors and labelled RNA or DNA probes of the fur gene and antibodies against furin can be used for diagnostic purposes.

(See also N70061, N70062, N70060, P70056 and P70058)
                                                                                                                                                                                                                                                                                                                                                                                            Identifying a modulator of intracellular signal transduction - I determining the interaction of a signal generating peptide with test substance, allows modulation of the immune system Example 8: Fig 15B: 74pp: English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N-PSDB; T60588.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-OCT-1996; U16195.
18-JUN-1996; US-665647.
10-OCT-1995; US-541964.
31-JAN-1996; US-594447.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  W15773;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W15773 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (TERR-) TERRAPIN TECHNOLOGIES INC.
Kauvar LM, Mochly-Rosen D, Napolitano EW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Recombinant DNA containing protein and antibodies and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Roenroek AJM,
Schalken JA;
WPI; 87-328946/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         36 CAQRCLQSCQQEPDDLKQKACESRCTKLEYDPRCV 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25 CHSECLGNCSQ-PDDPT-K-CVCACRNFYLDGRCV 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16.6%;
Similarity 42.9%;
                                                                          252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 149 AA;
 16.6%;
llarity 35.7%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the fur gene - used as a diagnostic aid
Score 75; DB 1;
Pred. No. 2.74e+01
6; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 75; DB 1;
Pred. No. 2.74e+01
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 10;
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54

104 NECIPSTSAELESEARSILEPDTSNSRSSSAFEAAKSROP 143

Mismatches

13;

Indels

2;

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2

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Query Match
Best Local S
Matches 1
                                                                                                                                                       and growth regulation or the organisation of the cytoskeleton. Polypeptides containing a WW domain are identified by treating a multivalent recognition unit complex that has selective binding affinity for a WW domain, with many polypeptides and identifying those with selective affinity for the complex. Proteins containing WW domains are used for targeted drug screening, i.e. to identify potential modulators of specific WW domain interactions. The valency of the recognition unit is important in determining specificity of interaction with WW domains. In multivalent form specificity is elaxed, but not lost, so proteins containing WW domains similar, but not identical, to the sequence of the peptides' target WW can be detected, including new polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents a novel protein WWP1. The WWP1 gene was identified and isolated from human bone marrow and brain cDNA libraries, using peptides W38103-05. These peptide recognition units are based on the sequences of WW domain binding domains of the YAP WW domain binding proteins WBP-1 and WBP-2. The WW domain is a small functional domain found in a large number of proteins from a variety of species including humans, nematodes and yeast. Its name is derived from the observation that two tryptophan residues, one in the amino terminal portion of the WW domain and one in the carboxyl terminal portion, are conserved. Most proteins containing WW domains have a function involving cell signalling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-APR-1996; US-630916.

(CYNTO-) CYTOGEN CORP.

(UYNC-) UNIV NORTH CAROLINA.

FOWLKES DM, KAY BK, PITOZZI

WPI: 97-503324/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel human protein, designated WWP1, which contains WW domains.

Peptide recognition unit; YAP WW domain binding protein; WBP-1; WBP-2;

WW domain; cell signalling; growth regulation; cytoskeleton organisation;

***roated drug screening: modulator; WW domain interaction; WWP1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         useful in targetted drug selection Claim 48; Fig 17; 220pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying cell signalling and growth regulatory polypeptides by reaction with multivalent recognition complex - polypeptides are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  W09737223-A1.
09-OCT-1997.
03-APR-1997; U05547.
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W36794;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB; T95697
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Domain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Domain
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Similarity
11; Conser
                                                                                                                                     target W
larity 27.5%;
Conservative
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267. .305
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307. .3
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192. .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              . 229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WW.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "WW domain 1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "WW domain
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Score 74;
Pred. No.
14; Misma
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                                  DB 1;
3.34e+01
                                                             Length 683
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